

# **Supplementary material for “Cultural differences in strength of conformity explained through pathogen stress: A statistical test using hierarchical Bayesian estimation”**

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## **Supplementary Figures**

Supplementary Figure S1–S15: Scatter plots displaying the correlations between regional level of government effectiveness (pathogen stress or GDP per capita) and each index of regional strength of conformity by each global region.

## **Supplementary Tables**

Supplementary Table S1: The number of countries or regions in each global region used for analysis.

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Supplementary Table S5: Numerical values of estimated parameters in Model 3 using both pathogen stress and government effectiveness as independent variables.

Supplementary Table S6: Summary of analysis results using both pathogen stress and government effectiveness as independent variables.

Supplementary Table S7: Numerical values of estimated parameters in Model 1 using both pathogen stress and GDP per capita as independent variables.

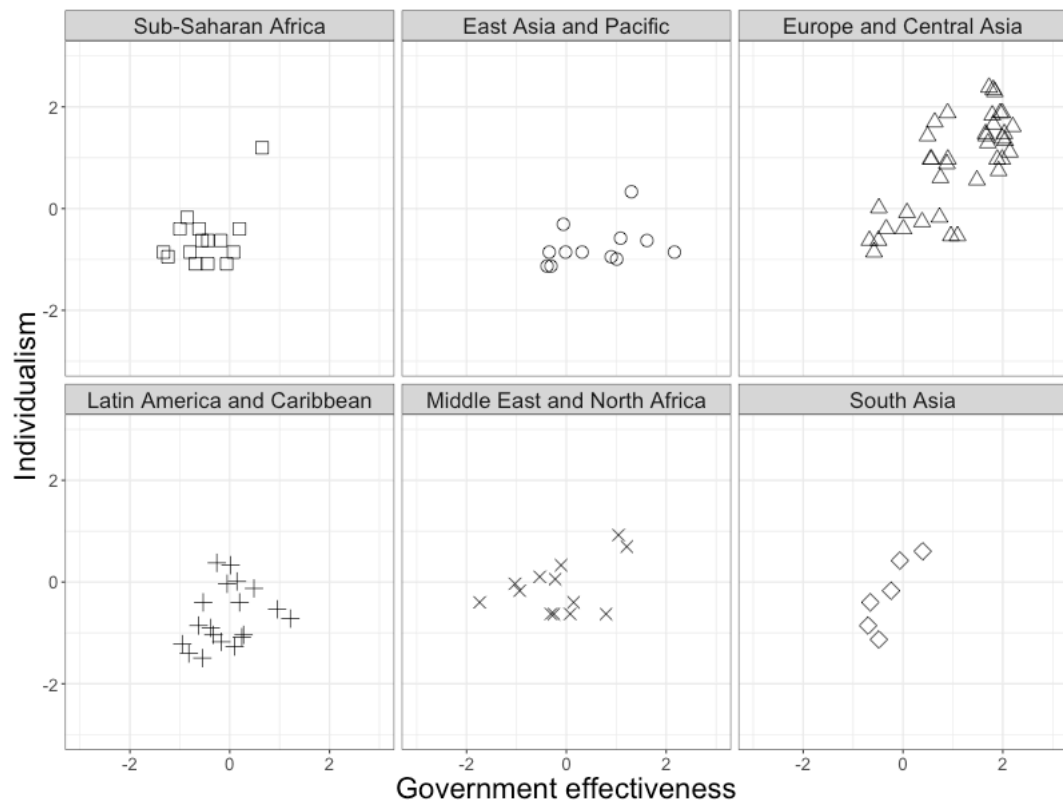
Supplementary Table S8: Numerical values of estimated parameters in Model 2 using both pathogen stress and GDP per capita as independent variables.

Supplementary Table S9: Numerical values of estimated parameters in Model 3 using both pathogen stress and GDP per capita as independent variables.

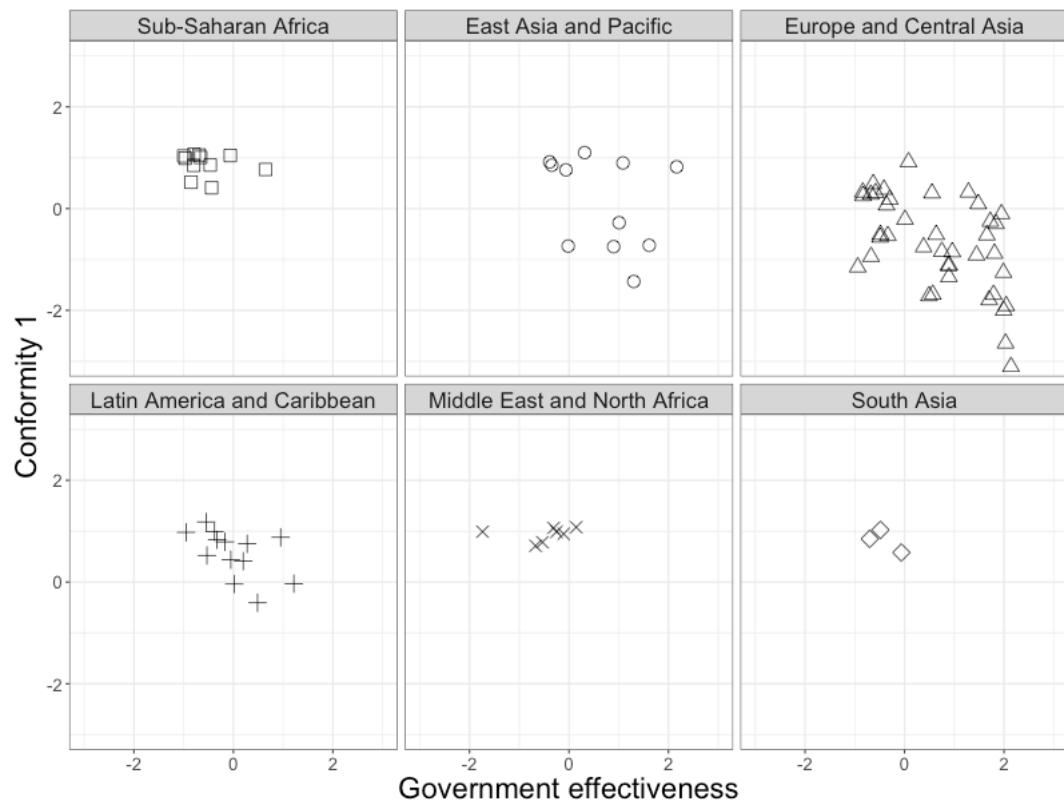
Supplementary Tables S10: Summary of analysis results using both pathogen stress and GDP per capita as independent variables.

## **Supplementary Method**

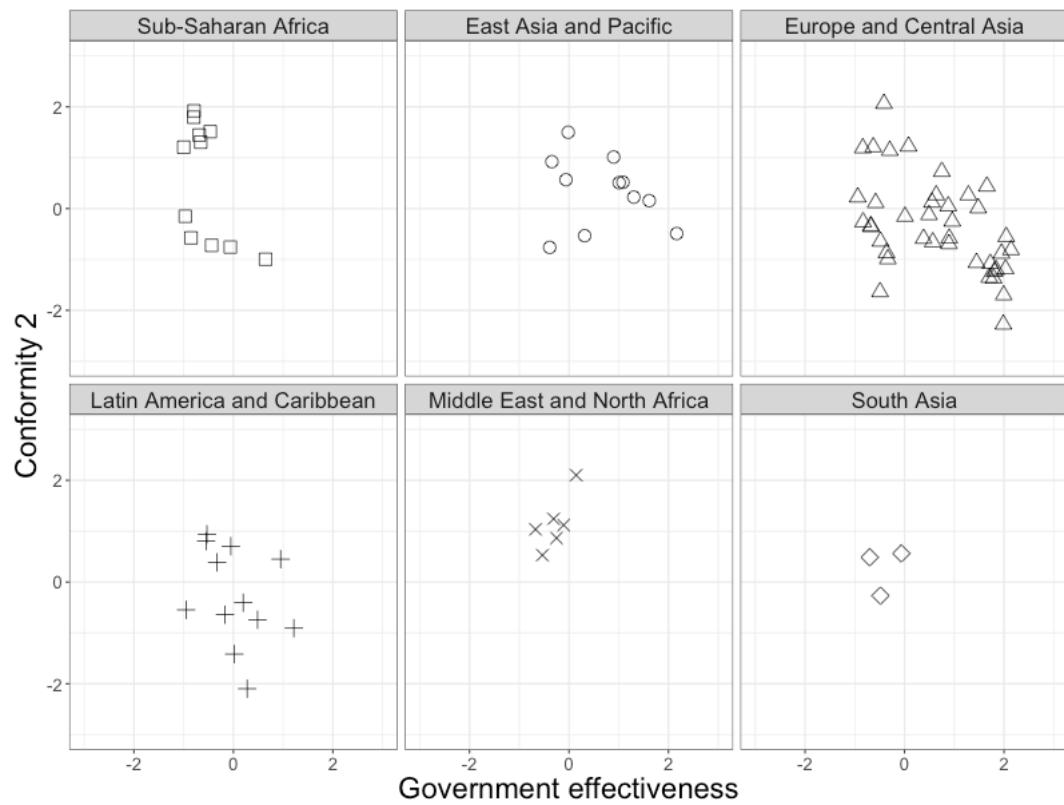
Bayesian estimation of zero-order correlation coefficients



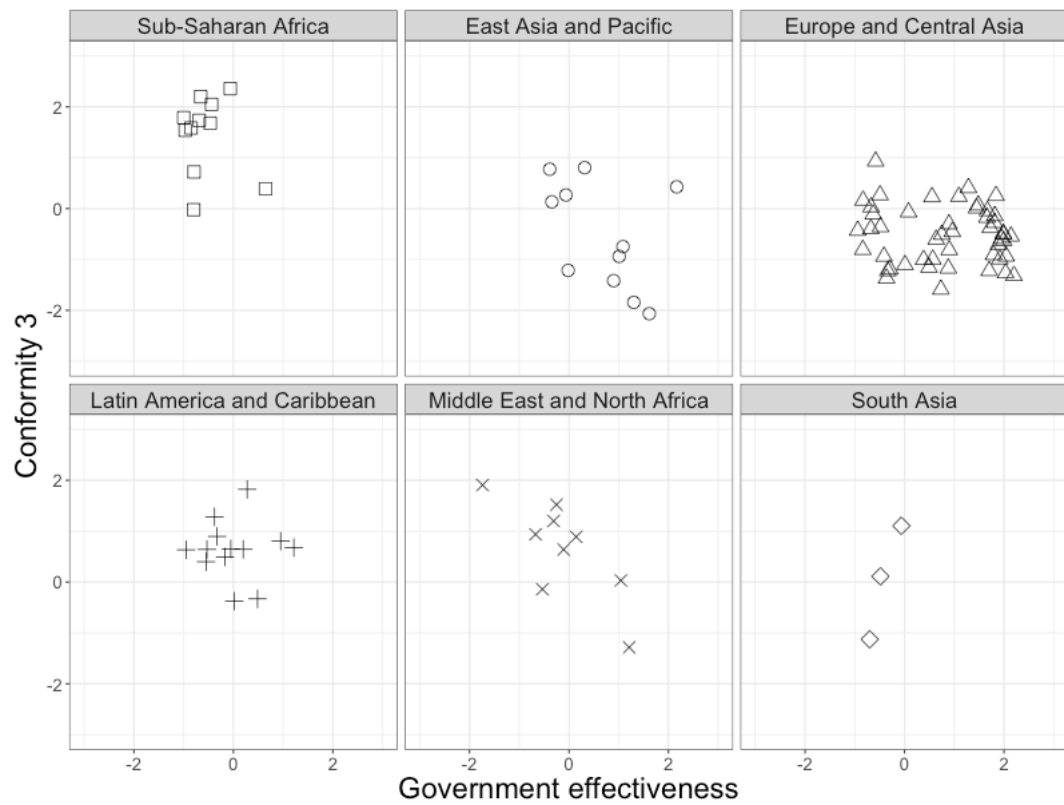
Supplementary Figure S1: Scatter plots displaying the correlation between regional level of government effectiveness and the indexes of *Individualism* by global region. Each point represents a country or region.



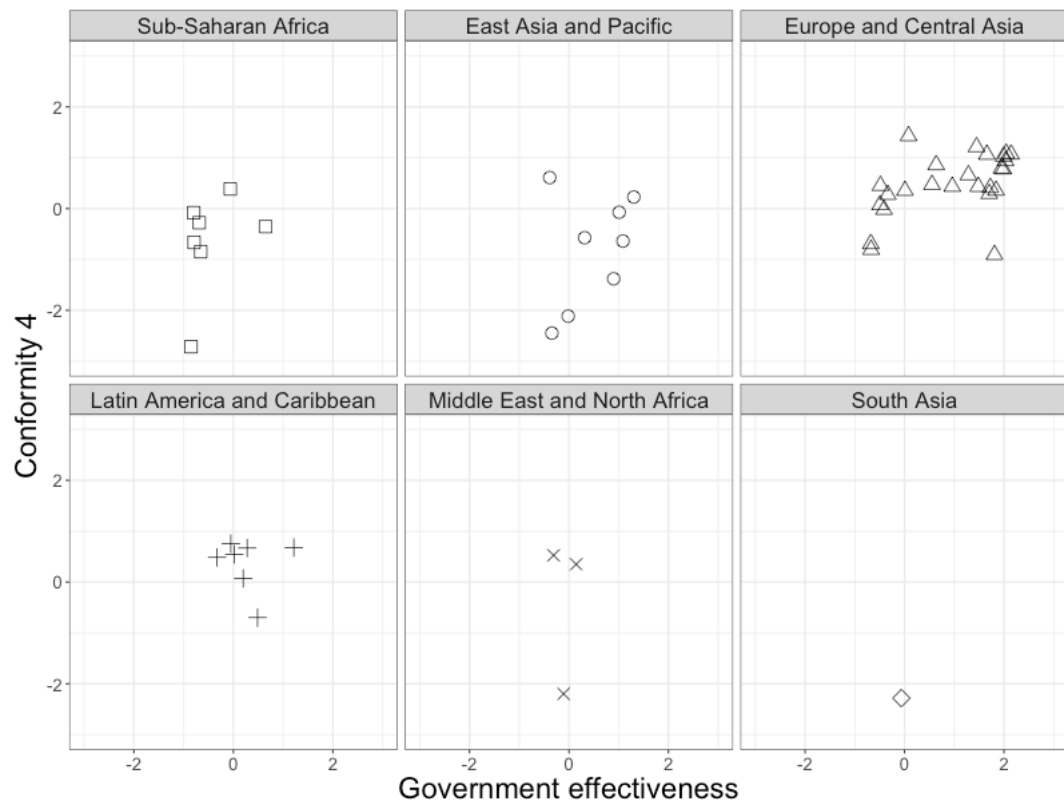
Supplementary Figure S2: Scatter plots displaying the correlation between regional level of government effectiveness and the indexes of *Conformity 1* by global region. Each point represents a country or region.



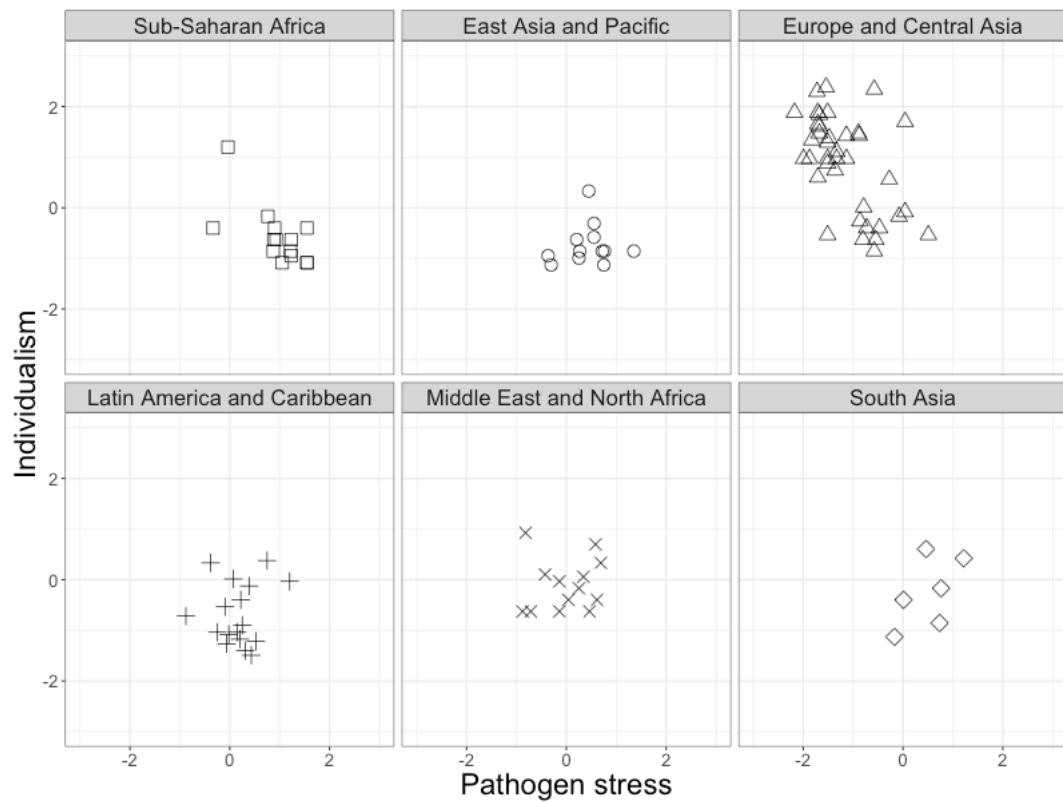
Supplementary Figure S3: Scatter plots displaying the correlation between regional level of government effectiveness and the indexes of *Conformity 2* by global region. Each point represents a country or region.



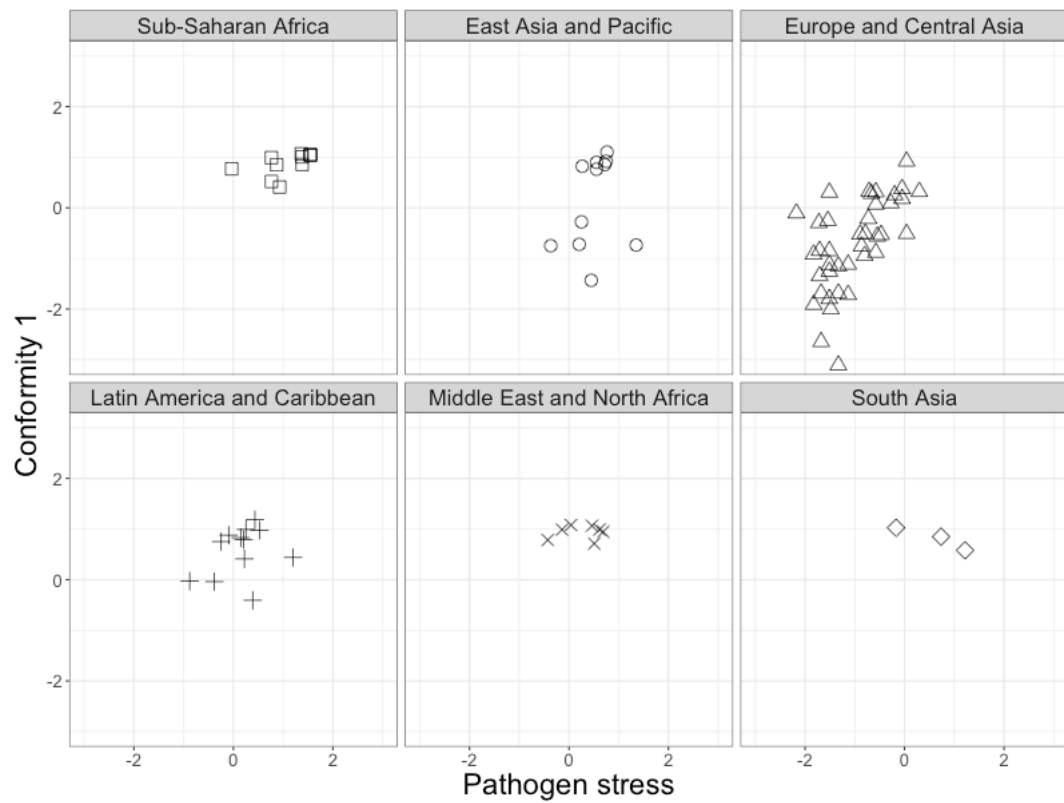
Supplementary Figure S4: Scatter plots displaying the correlation between regional level of government effectiveness and the indexes of *Conformity 3* by global region. Each point represents a country or region.



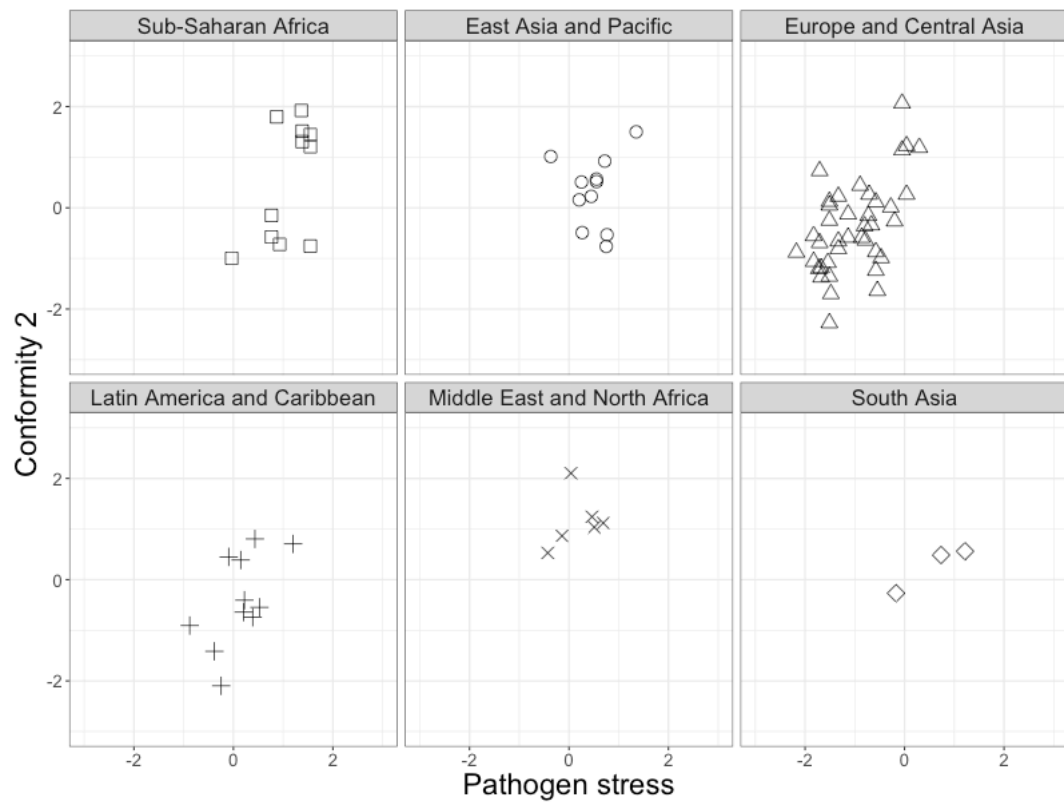
Supplementary Figure S5: Scatter plots displaying the correlation between regional level of government effectiveness and the indexes of *Conformity 4* by global region. Each point represents a country or region.



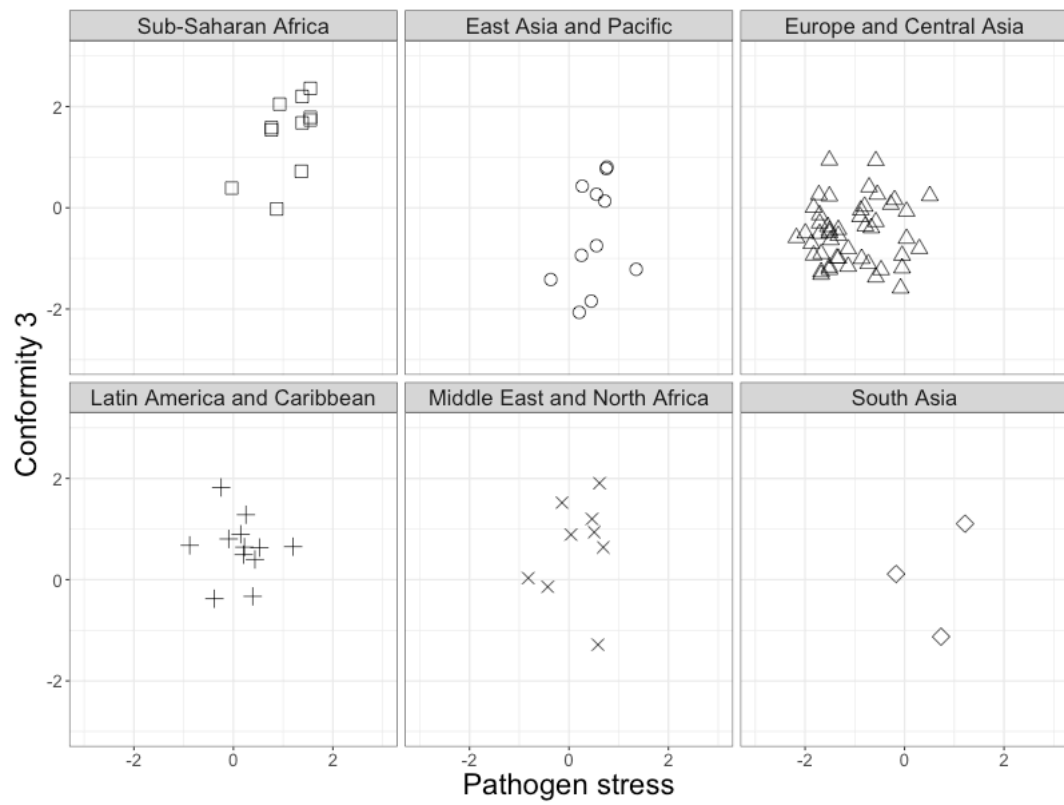
Supplementary Figure S6: Scatter plots displaying the correlation between regional level of pathogen stress and the index of *Individualism* by each global region. Each point represents a country or region.



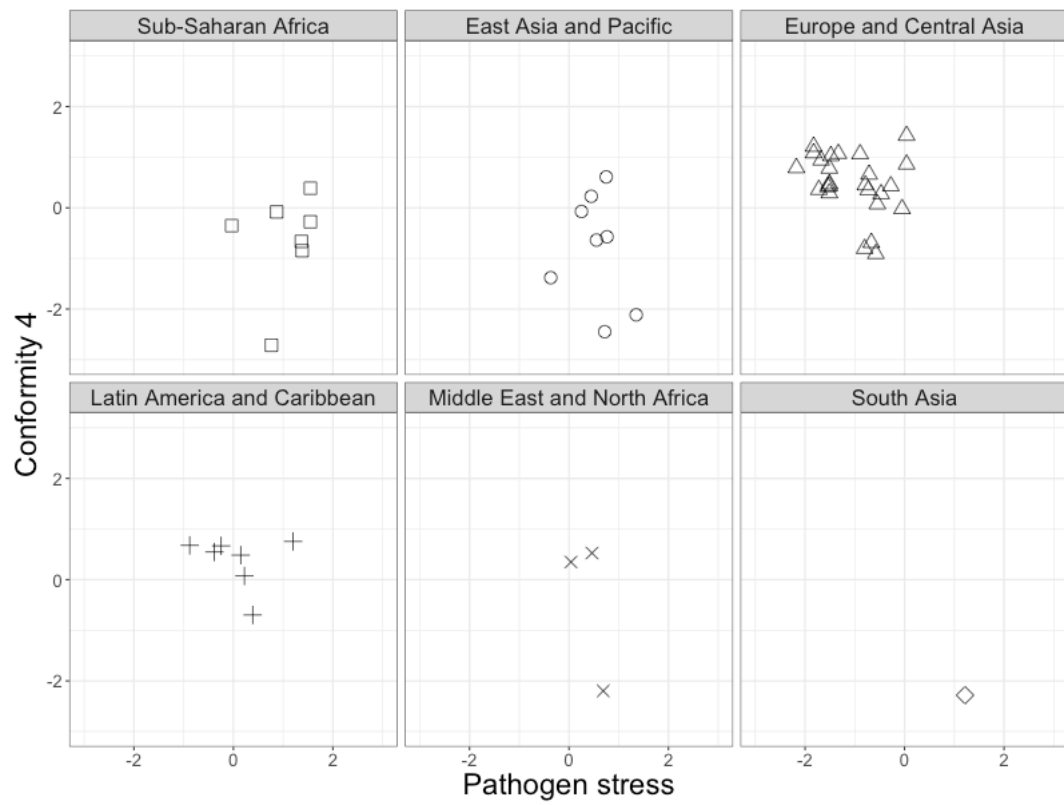
Supplementary Figure S7: Scatter plots displaying the correlation between regional level of pathogen stress and the index of *Conformity 1* by each global region. Each point represents a country or region.



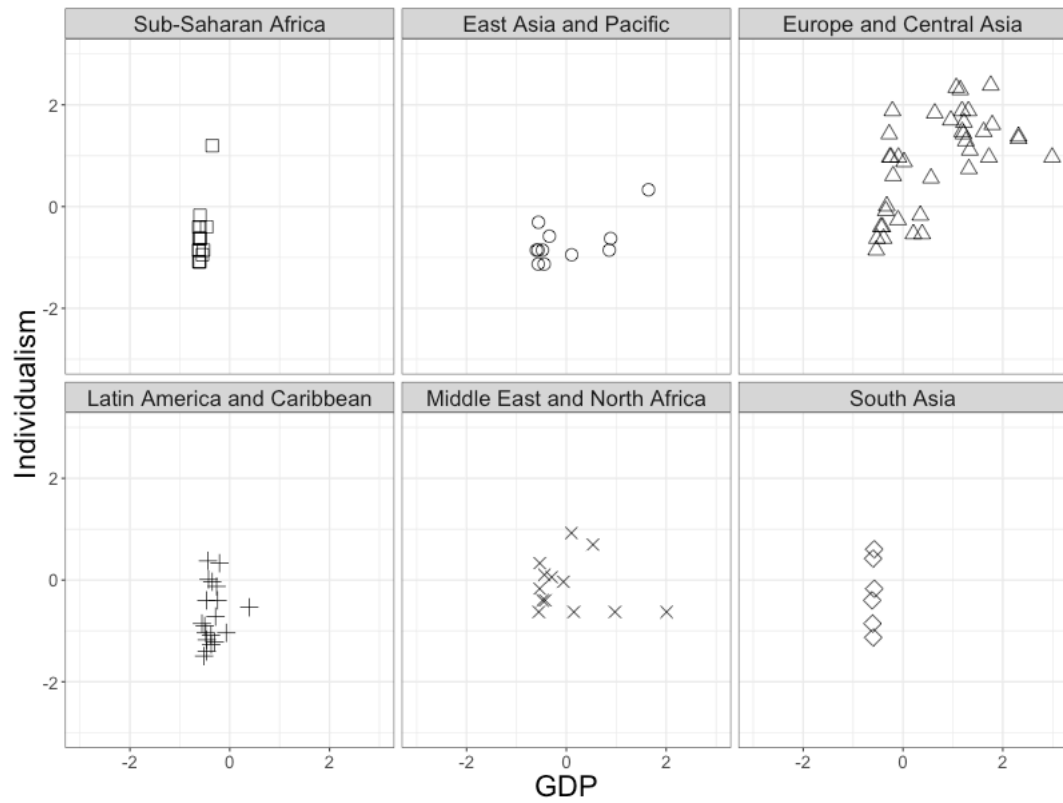
Supplementary Figure S8: Scatter plots displaying the correlation between regional level of pathogen stress and the index of *Conformity 2* by global region. Each point represents a country or region.



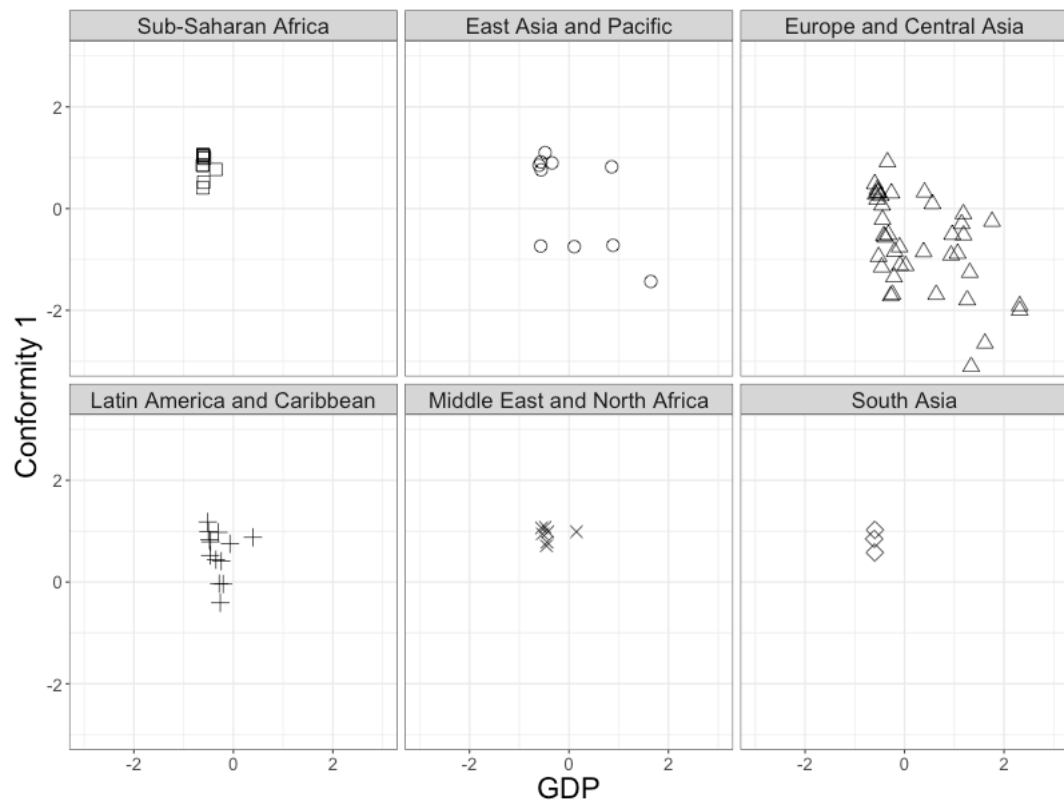
Supplementary Figure S9: Scatter plots displaying the correlation between regional level of pathogen stress and the index of *Conformity 3* by global region. Each point represents a country or region.



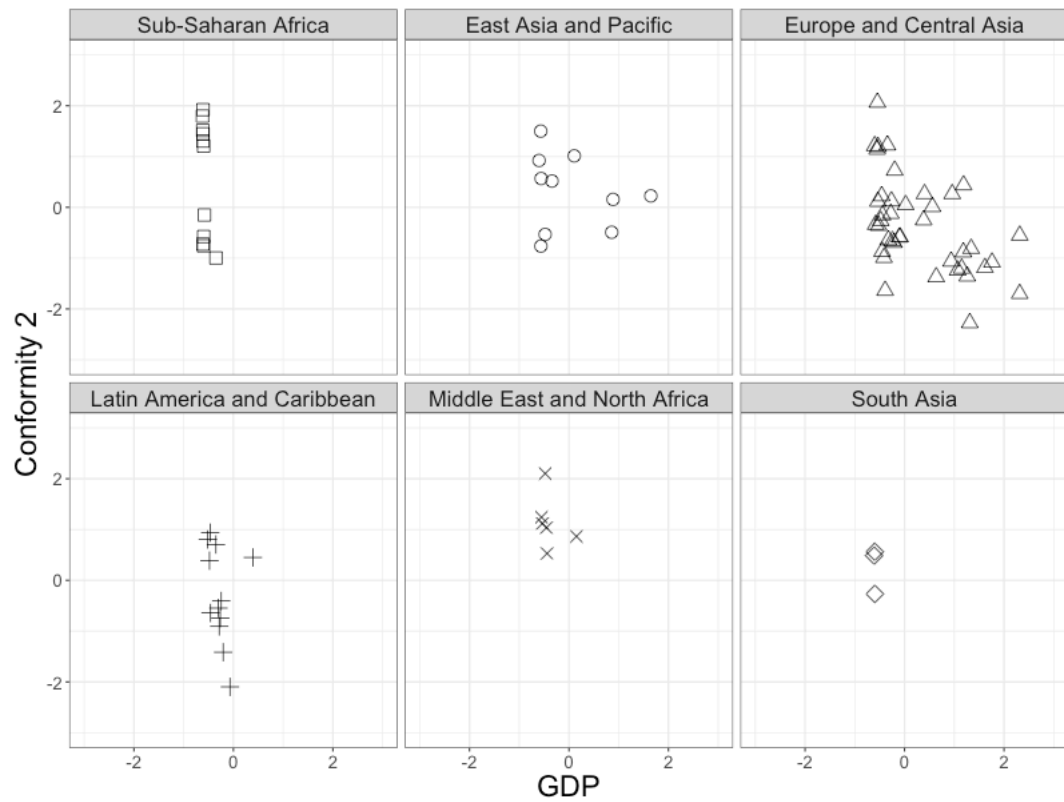
Supplementary Figure S10: Scatter plots displaying a correlation between regional level of pathogen stress and the index of *Conformity 4* by global region. Each point represents a country or region.



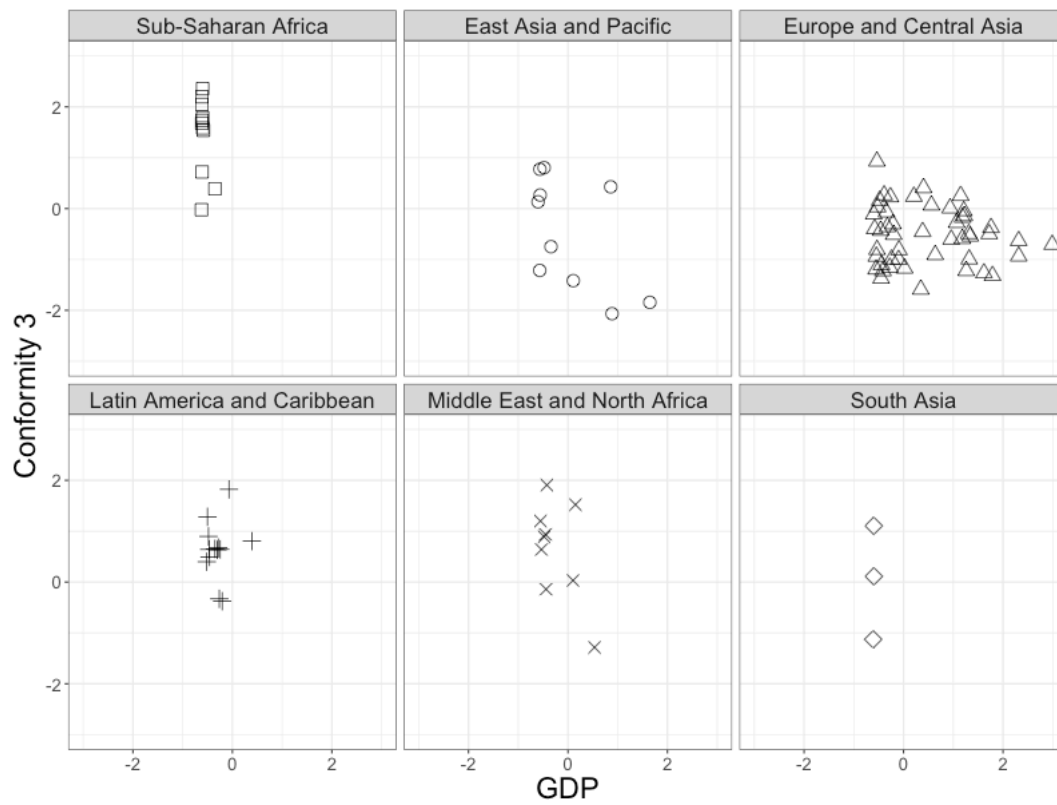
Supplementary Figure S11: Scatter plots displaying the correlation between regional level of GDP per capita and the index of *Individualism* by global region. Each point represents a country or region.



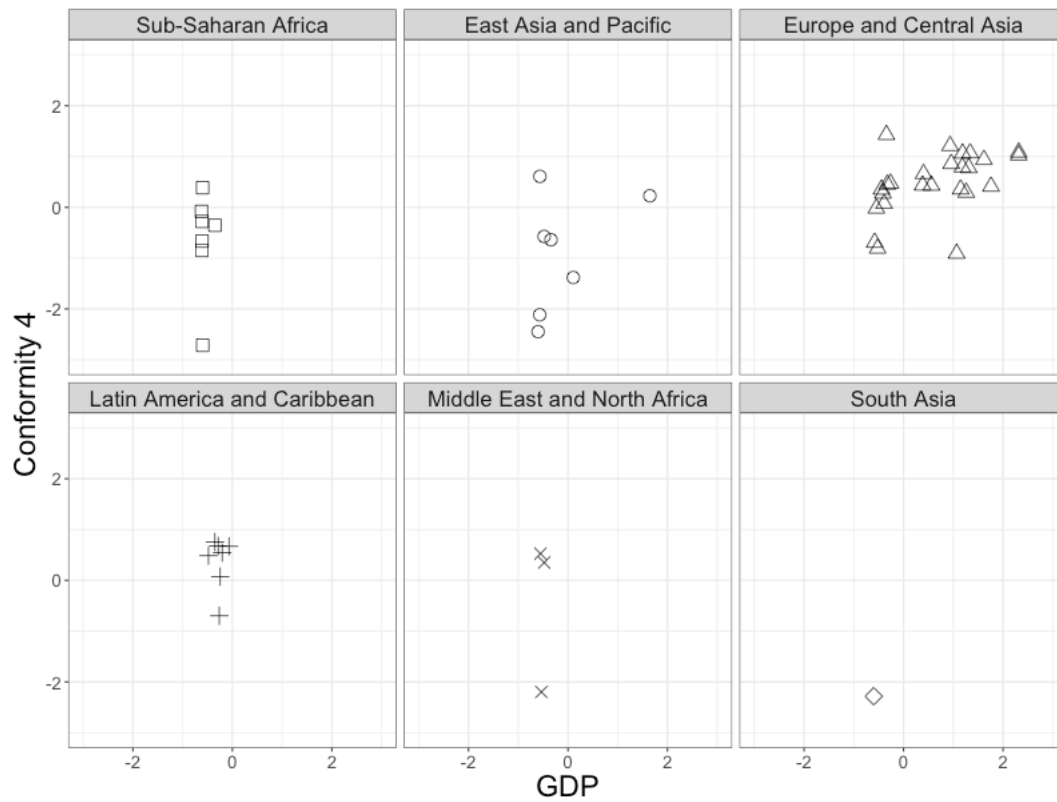
Supplementary Figure S12: Scatter plots displaying the correlation between regional level of GDP per capita and the index of *Conformity 1* by global region. Each point represents a country or region.



Supplementary Figure S13: Scatter plots displaying the correlation between regional level of GDP per capita and the index of *Conformity 2* by global region. Each point represents a country or region.



Supplementary Figure S14: Scatter plots displaying the correlation between regional level of GDP per capita and the index of *Conformity 3* by global region. Each point represents a country or region.



Supplementary Figure S15: Scatter plots displaying the correlation between regional level of GDP per capita and the index of *Conformity 4* by global region. Each point represents a country or region.

Supplementary Table S1: The number of countries or regions in each global region used for analysis (GE: government effectiveness, PS: pathogen stress, IND: *Individualism*, C1: *Conformity 1*, C2: *Conformity 2*, C3, *Conformity 3*, C4: *Conformity 4*).

	Dependent variables				
	IND	C1	C2	C3	C4
GE and PS were used as independent variables					
Sub-Saharan Africa	14	11	11	11	7
East Asia and Pacific	12	11	11	11	8
Europe and Central Asia	38	39	39	47	25
Latin America and Caribbean	17	12	11	12	7
Middle East and North Africa	13	7	6	9	3
South Asia	6	3	3	3	1
Total	100	83	81	93	51
GDP per capita and PS were used for independent variables					
Sub-Saharan Africa	14	11	11	11	7
East Asia and Pacific	11	10	10	10	7
Europe and Central Asia	38	39	39	47	25
Latin America and Caribbean	17	12	11	12	7
Middle East and North Africa	13	7	6	9	3
South Asia	6	3	3	3	1
Total	99	82	80	92	50

Supplementary Table S2: Posterior distribution of zero-order correlation coefficients between dependent variables. *ESS* represents effective sample sizes of MCMC simulations.

	<i>Mean</i>	<i>SD</i>	Quantiles			Sample size	<i>ESS</i>
			2.5%	50%	97.5%		
<i>Individualism and Conformity 1</i>	−0.62	0.08	−0.75	−0.62	−0.45	70	12142
<i>Individualism and Conformity 2</i>	−0.40	0.10	−0.59	−0.40	−0.18	68	14609
<i>Individualism and Conformity 3</i>	−0.40	0.09	−0.58	−0.41	−0.21	80	14885
<i>Individualism and Conformity 4</i>	0.27	0.14	−0.03	0.27	0.52	45	16000
<i>Conformity 1 and Conformity 2</i>	0.47	0.09	0.29	0.47	0.63	83	16000
<i>Conformity 1 and Conformity 3</i>	0.66	0.06	0.53	0.66	0.77	85	10564
<i>Conformity 1 and Conformity 4</i>	−0.30	0.13	−0.53	−0.31	−0.04	51	16000
<i>Conformity 2 and Conformity 3</i>	0.15	0.11	−0.06	0.15	0.36	83	16000
<i>Conformity 2 and Conformity 4</i>	−0.32	0.13	−0.55	−0.32	−0.05	51	16000
<i>Conformity 3 and Conformity 4</i>	−0.16	0.14	−0.42	−0.16	0.12	51	16000

Supplementary Table S3: Numerical values of estimated parameter in Model 1 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Individualism						
$a_0$	−0.22	0.07	−0.37	−0.22	−0.08	13857
GE	0.44	0.10	0.25	0.44	0.63	10736
PS	−0.39	0.10	−0.58	−0.39	−0.19	11323
$\sigma_y$	0.69	0.05	0.60	0.69	0.80	13446
Conformity 1						
$a_0$	0.19	0.08	0.04	0.19	0.34	16000
GE	−0.31	0.10	−0.50	−0.31	−0.12	12934
PS	0.55	0.10	0.36	0.55	0.74	13420
$\sigma_y$	0.68	0.05	0.58	0.67	0.79	14347
Conformity 2						
$a_0$	0.13	0.10	−0.06	0.13	0.33	14358
GE	−0.20	0.12	−0.45	−0.20	0.04	11178
PS	0.43	0.12	0.20	0.43	0.67	11891
$\sigma_y$	0.84	0.07	0.72	0.84	0.99	14594
Conformity 3						
$a_0$	0.19	0.10	0.002	0.19	0.39	15191
GE	−0.25	0.11	−0.47	−0.25	−0.02	11100
PS	0.36	0.12	0.13	0.36	0.59	11223
$\sigma_y$	0.86	0.06	0.74	0.85	0.99	13363
Conformity 4						
$a_0$	−0.17	0.14	−0.45	−0.17	0.11	9963
GE	0.15	0.18	−0.21	0.15	0.51	7762
PS	−0.42	0.17	−0.76	−0.42	−0.09	8812
$\sigma_y$	0.88	0.09	0.72	0.87	1.09	11771

Supplementary Table S4: Numerical values of estimated parameter in Model 2 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$  represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia).  $ESS$  represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Individualism						
$\mu_a$	−0.29	0.29	−0.87	−0.29	0.29	8551
GE	0.54	0.09	0.35	0.54	0.72	16000
PS	−0.11	0.12	−0.35	−0.11	0.13	10850
$a_1$	−0.20	0.18	−0.55	−0.19	0.16	16000
$a_2$	−0.93	0.20	−1.31	−0.93	−0.52	12602
$a_3$	0.14	0.15	−0.16	0.14	0.45	14116
$a_4$	−0.64	0.15	−0.93	−0.64	−0.36	16000
$a_5$	−0.06	0.16	−0.38	−0.06	0.26	16000
$a_6$	−0.10	0.23	−0.54	−0.09	0.36	16000
$\sigma_a$	0.61	0.33	0.25	0.53	1.44	7018
$\sigma_y$	0.60	0.05	0.52	0.60	0.70	16000
Conformity 1						
$\mu_a$	0.28	0.21	−0.09	0.27	0.72	5287
GE	−0.30	0.10	−0.49	−0.30	−0.10	11251
PS	0.43	0.14	0.14	0.43	0.70	7365
$a_1$	0.26	0.20	−0.11	0.25	0.66	10565
$a_2$	0.17	0.19	−0.21	0.17	0.55	12158
$a_3$	−0.03	0.16	−0.36	−0.03	0.26	6655
$a_4$	0.44	0.18	0.10	0.44	0.80	7078
$a_5$	0.51	0.23	0.09	0.50	0.99	7103
$a_6$	0.35	0.27	−0.15	0.33	0.94	9989
$\sigma_a$	0.37	0.26	0.08	0.31	1.02	3637
$\sigma_y$	0.65	0.05	0.55	0.65	0.77	11364

Supplementary Table S4 (cond.): Numerical values of estimated parameters in Model 2 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$  represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

			Quantiles			
	Mean	SD	2.5%	50%	97.5%	ESS
Conformity 2						
$\mu_a$	0.11	0.32	−0.54	0.11	0.74	9305
GE	−0.17	0.13	−0.42	−0.17	0.08	5108
PS	0.58	0.18	0.25	0.58	0.94	7998
$a_1$	−0.13	0.27	−0.66	−0.12	0.36	3075
$a_2$	0.15	0.25	−0.36	0.15	0.65	12278
$a_3$	0.31	0.18	−0.05	0.31	0.68	10656
$a_4$	−0.33	0.25	−0.82	−0.34	0.17	1164
$a_5$	0.71	0.33	0.12	0.71	1.37	4165
$a_6$	−0.04	0.37	−0.79	−0.02	0.66	16000
$\sigma_a$	0.62	0.40	0.14	0.53	1.62	3274
$\sigma_y$	0.79	0.07	0.67	0.78	0.96	616
Conformity 3						
$\mu_a$	0.27	0.43	−0.59	0.27	1.16	9214
GE	−0.14	0.11	−0.35	−0.14	0.07	13657
PS	0.12	0.15	−0.17	0.12	0.42	10529
$a_1$	1.16	0.27	0.65	1.16	1.69	12857
$a_2$	−0.42	0.25	−0.91	−0.43	0.06	13922
$a_3$	−0.27	0.17	−0.60	−0.27	0.05	12949
$a_4$	0.60	0.20	0.19	0.60	0.99	16000
$a_5$	0.56	0.23	0.10	0.56	1.02	16000
$a_6$	−0.01	0.38	−0.75	−0.01	0.72	16000
$\sigma_a$	0.91	0.49	0.38	0.79	2.15	7100
$\sigma_y$	0.73	0.06	0.62	0.72	0.85	16000

Supplementary Table S4 (cond.): Numerical values of estimated parameters in Model 2 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$  represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 4						
$\mu_a$	−0.39	0.45	−1.39	−0.34	0.34	5231
GE	0.27	0.18	−0.10	0.27	0.62	6319
PS	−0.16	0.25	−0.63	−0.16	0.35	3443
$a_1$	−0.37	0.34	−1.08	−0.35	0.25	5874
$a_2$	−0.71	0.37	−1.44	−0.72	−0.03	3103
$a_3$	0.03	0.25	−0.43	0.02	0.54	4397
$a_4$	0.12	0.30	−0.43	0.11	0.73	5725
$a_5$	−0.35	0.39	−1.17	−0.33	0.41	9420
$a_6$	−1.06	0.79	−2.83	−0.95	0.12	3078
$\sigma_a$	0.75	0.57	0.07	0.63	2.17	2689
$\sigma_y$	0.82	0.09	0.66	0.81	1.03	6541

Supplementary Table S5: Numerical values of estimated parameters in Model 3 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$ , GE, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Individualism						
$\mu_a$	−0.38	0.26	−0.89	−0.38	0.13	6822
$\mu_{GE}$	0.48	0.20	0.09	0.47	0.87	6604
$\mu_{PS}$	0.07	0.27	−0.41	0.05	0.64	6929
$a_1$	−0.18	0.24	−0.64	−0.18	0.33	8870
$a_2$	−0.84	0.25	−1.34	−0.83	−0.37	5778
$a_3$	−0.17	0.18	−0.51	−0.17	0.18	10043
$a_4$	−0.70	0.15	−0.99	−0.70	−0.42	10297
$a_5$	−0.10	0.15	−0.39	−0.10	0.20	12569
$a_6$	−0.25	0.27	−0.78	−0.25	0.29	7768
GE <sub>1</sub>	0.45	0.24	−0.03	0.46	0.93	10631
GE <sub>2</sub>	0.25	0.20	−0.15	0.25	0.60	5397
GE <sub>3</sub>	0.66	0.12	0.42	0.66	0.91	5398
GE <sub>4</sub>	0.49	0.22	0.07	0.49	0.94	9425
GE <sub>5</sub>	0.36	0.18	−0.002	0.37	0.69	9464
GE <sub>6</sub>	0.64	0.34	0.08	0.59	1.45	7004
PS <sub>1</sub>	−0.19	0.25	−0.71	−0.17	0.27	8240
PS <sub>2</sub>	0.07	0.29	−0.47	0.06	0.66	8117
PS <sub>3</sub>	−0.27	0.17	−0.60	−0.27	0.05	5718
PS <sub>4</sub>	0.31	0.31	−0.23	0.30	0.95	6336
PS <sub>5</sub>	0.19	0.26	−0.30	0.18	0.73	8485
PS <sub>6</sub>	0.29	0.36	−0.32	0.26	1.05	6483
$\sigma_a$	0.51	0.30	0.16	0.44	1.27	5450
$\sigma_{GE}$	0.33	0.26	0.04	0.28	0.96	3146
$\sigma_{PS}$	0.46	0.32	0.07	0.40	1.28	4227
$\sigma_y$	0.56	0.04	0.48	0.56	0.65	16000

Supplementary Table S5 (cond.): Numerical values of estimated parameters in Model 3 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$ , GE, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	<i>Mean</i>	<i>SD</i>	Quantiles			<i>ESS</i>
			2.5%	50%	97.5%	
<i>Conformity 1</i>						
$\mu_a$	0.37	0.22	−0.10	0.36	0.82	953
$\mu_{\text{GE}}$	−0.27	0.17	−0.59	−0.27	0.08	5800
$\mu_{\text{PS}}$	0.33	0.24	−0.19	0.35	0.77	4562
$a_1$	0.40	0.26	−0.07	0.39	0.96	5750
$a_2$	0.25	0.23	−0.21	0.26	0.71	4975
$a_3$	0.10	0.19	−0.29	0.10	0.47	1382
$a_4$	0.48	0.18	0.14	0.47	0.83	6079
$a_5$	0.57	0.25	0.14	0.55	1.11	3353
$a_6$	0.44	0.29	−0.09	0.42	1.10	5386
GE <sub>1</sub>	−0.22	0.22	−0.61	−0.24	0.31	5848
GE <sub>2</sub>	−0.29	0.16	−0.63	−0.29	0.03	7327
GE <sub>3</sub>	−0.27	0.11	−0.50	−0.27	−0.05	6332
GE <sub>4</sub>	−0.28	0.20	−0.69	−0.28	0.13	7253
GE <sub>5</sub>	−0.24	0.21	−0.64	−0.26	0.24	5460
GE <sub>6</sub>	−0.29	0.28	−0.93	−0.29	0.26	6773
PS <sub>1</sub>	0.31	0.24	−0.20	0.32	0.74	4450
PS <sub>2</sub>	0.26	0.29	−0.38	0.29	0.78	4557
PS <sub>3</sub>	0.57	0.19	0.21	0.57	0.96	1976
PS <sub>4</sub>	0.26	0.29	−0.39	0.29	0.76	4466
PS <sub>5</sub>	0.31	0.34	−0.46	0.34	0.95	4719
PS <sub>6</sub>	0.27	0.33	−0.50	0.31	0.83	4412
$\sigma_a$	0.35	0.26	0.03	0.30	0.97	1821
$\sigma_{\text{GE}}$	0.19	0.20	0.01	0.13	0.71	1841
$\sigma_{\text{PS}}$	0.32	0.27	0.02	0.27	1.00	1933
$\sigma_y$	0.65	0.05	0.55	0.65	0.77	7140

Supplementary Table 5 (cond.): Numerical values of estimated parameter in Model 3 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$ , GE, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 2						
$\mu_a$	0.08	0.37	−0.67	0.09	0.78	4666
$\mu_{GE}$	−0.17	0.26	−0.67	−0.17	0.36	5703
$\mu_{PS}$	0.58	0.27	0.08	0.59	1.10	1551
$a_1$	−0.29	0.39	−1.13	−0.26	0.39	3265
$a_2$	0.15	0.31	−0.47	0.15	0.79	3437
$a_3$	0.30	0.21	−0.12	0.30	0.71	3348
$a_4$	−0.38	0.25	−0.87	−0.38	0.12	2774
$a_5$	0.76	0.36	0.08	0.75	1.46	3661
$a_6$	−0.05	0.42	−0.91	−0.04	0.75	3457
$GE_1$	−0.34	0.36	−1.30	−0.28	0.22	941
$GE_2$	−0.11	0.22	−0.52	−0.12	0.37	4179
$GE_3$	−0.19	0.14	−0.46	−0.18	0.09	4063
$GE_4$	−0.11	0.28	−0.63	−0.12	0.50	4815
$GE_5$	−0.13	0.43	−1.04	−0.15	0.89	797
$GE_6$	−0.14	0.41	−0.97	−0.15	0.76	7342
$PS_1$	0.65	0.30	0.10	0.64	1.33	2069
$PS_2$	0.47	0.34	−0.32	0.50	1.07	2373
$PS_3$	0.56	0.21	0.14	0.56	0.96	1813
$PS_4$	0.66	0.32	0.08	0.64	1.35	1729
$PS_5$	0.59	0.37	−0.17	0.59	1.35	2362
$PS_6$	0.58	0.34	−0.12	0.58	1.28	2852
$\sigma_a$	0.69	0.46	0.16	0.60	1.78	2550
$\sigma_{GE}$	0.33	0.33	0.03	0.23	1.16	1344
$\sigma_{PS}$	0.31	0.29	0.03	0.23	1.06	1589
$\sigma_y$	0.79	0.07	0.67	0.78	0.93	7190

Supplementary Table S5 (cond.): Numerical values of estimated parameter in Model 3 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$ , GE, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 3						
$\mu_a$	0.21	0.38	−0.53	0.21	0.95	4414
$\mu_{\text{GE}}$	−0.17	0.39	−0.83	−0.20	0.69	4868
$\mu_{\text{PS}}$	0.21	0.29	−0.35	0.21	0.77	5613
$a_1$	0.71	0.41	−0.10	0.72	1.50	4360
$a_2$	−0.23	0.33	−0.87	−0.23	0.39	4917
$a_3$	−0.35	0.18	−0.71	−0.36	0.004	4583
$a_4$	0.58	0.20	0.19	0.58	0.97	5160
$a_5$	0.47	0.23	0.04	0.47	0.92	3679
$a_6$	0.10	0.46	−0.77	0.08	1.08	4360
GE <sub>1</sub>	−0.15	0.38	−0.88	−0.17	0.66	3778
GE <sub>2</sub>	−0.47	0.25	−0.97	−0.46	0.02	3841
GE <sub>3</sub>	−0.03	0.12	−0.26	−0.02	0.22	2877
GE <sub>4</sub>	−0.07	0.32	−0.67	−0.08	0.59	6056
GE <sub>5</sub>	−0.64	0.28	−1.22	−0.63	−0.13	3709
GE <sub>6</sub>	0.36	0.81	−0.74	0.16	2.45	3291
PS <sub>1</sub>	0.55	0.38	−0.08	0.50	1.38	3004
PS <sub>2</sub>	0.16	0.34	−0.54	0.16	0.83	6592
PS <sub>3</sub>	0.13	0.18	−0.21	0.13	0.48	3502
PS <sub>4</sub>	0.07	0.33	−0.65	0.10	0.69	5464
PS <sub>5</sub>	0.13	0.32	−0.56	0.14	0.73	6627
PS <sub>6</sub>	0.23	0.36	−0.51	0.22	0.99	8003
$\sigma_a$	0.73	0.43	0.25	0.63	1.78	3189
$\sigma_{\text{GE}}$	0.65	0.56	0.12	0.51	2.07	3254
$\sigma_{\text{PS}}$	0.42	0.37	0.06	0.33	1.29	2123
$\sigma_y$	0.69	0.06	0.59	0.69	0.82	1670

Supplementary Table S5 (cond.): Numerical values of estimated parameter in Model 3 using both pathogen stress and government effectiveness as independent variables. GE and PS represent government effectiveness and pathogen stress, respectively. Subscript numbers under  $a$ , GE, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	<i>Mean</i>	<i>SD</i>	Quantiles			<i>ESS</i>
			2.5%	50%	97.5%	
<i>Conformity 4</i>						
$\mu_a$	−0.29	0.47	−1.36	−0.24	0.49	4546
$\mu_{GE}$	0.30	0.51	−0.63	0.29	1.27	6272
$\mu_{PS}$	−0.34	0.52	−1.53	−0.29	0.53	3730
$a_1$	−0.42	0.47	−1.47	−0.37	0.38	4220
$a_2$	−0.61	0.48	−1.58	−0.59	0.24	2711
$a_3$	0.07	0.27	−0.44	0.06	0.62	6340
$a_4$	0.16	0.31	−0.42	0.15	0.79	5494
$a_5$	−0.19	0.45	−1.10	−0.18	0.72	4996
$a_6$	−0.75	0.87	−2.76	−0.58	0.57	2963
$GE_1$	0.50	0.48	−0.32	0.43	1.64	4225
$GE_2$	0.29	0.36	−0.45	0.28	1.03	5584
$GE_3$	0.27	0.20	−0.13	0.27	0.64	6923
$GE_4$	0.19	0.47	−0.88	0.22	1.09	7413
$GE_5$	0.21	0.80	−1.70	0.26	1.68	6341
$GE_6$	0.36	1.05	−1.38	0.30	2.44	4022
$PS_1$	0.04	0.46	−0.76	0.00	1.07	3515
$PS_2$	−0.37	0.51	−1.47	−0.34	0.57	3217
$PS_3$	−0.12	0.29	−0.68	−0.12	0.47	5308
$PS_4$	−0.15	0.42	−0.99	−0.15	0.70	7933
$PS_5$	−0.69	0.88	−2.88	−0.49	0.57	2991
$PS_6$	−0.71	0.77	−2.46	−0.57	0.49	2630
$\sigma_a$	0.72	0.61	0.05	0.58	2.27	2350
$\sigma_{GE}$	0.57	0.74	0.03	0.36	2.34	1372
$\sigma_{PS}$	0.69	0.69	0.04	0.51	2.39	2082
$\sigma_y$	0.82	0.10	0.66	0.81	1.03	9234

Supplementary Tables S6: Summary of analysis results using both pathogen stress and government effectiveness as independent variables. Parameters in parentheses indicate the parameters in Model 3. Asterisks in columns of global effects (GE, PS,  $\mu_{GE}$ , or  $\mu_{PS}$ ) indicate that the effect was significant (i.e., the 95% Bayesian credible interval of the effect did not include zero). Numbers in columns of region-specific effects ( $GE_j$  or  $PS_j$ ) indicate global regions in which significant effects were found (1 = Sub-Saharan Africa, 2 = East Asia and Pacific, 3 = Europe and Central Asia, 4 = Latin America and Caribbean, 5 = Middle East and North Africa, and 6 = South Asia). Rows shaded in gray indicate the best model, in which the WAIC value was smallest in three models.

Dependent variables	Model	Parameters			
		GE ( $\mu_{GE}$ )	PS ( $\mu_{PS}$ )	( $GE_j$ )	( $PS_j$ )
<i>Individualism</i>	Model 1	*	*		
	Model 2	*			
	Model 3	*		3,4,6	
<i>Conformity 1</i>	Model 1	*	*		
	Model 2	*	*		
	Model 3			3	3
<i>Conformity 2</i>	Model 1		*		
	Model 2		*		
	Model 3		*		1, 3, 4
<i>Conformity 3</i>	Model 1	*	*		
	Model 2				
	Model 3			5	
<i>Conformity 4</i>	Model 1		*		
	Model 2				
	Model 3				

Supplementary Table S7: Numerical values of estimated parameters in Model 1 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

			Quantiles			
	Mean	SD	2.5%	50%	97.5%	ESS
Individualism						
$a_0$	−0.09	0.07	−0.24	−0.09	0.05	15682
GDP	0.40	0.11	0.18	0.40	0.62	12681
PS	−0.46	0.10	−0.66	−0.46	−0.27	12240
$\sigma_y$	0.72	0.05	0.63	0.72	0.83	13678
Conformity 1						
$a_0$	0.09	0.08	−0.06	0.09	0.25	12862
GDP	−0.45	0.12	−0.69	−0.45	−0.22	13024
PS	0.54	0.09	0.36	0.54	0.72	12445
$\sigma_y$	0.66	0.05	0.57	0.66	0.78	13877
Conformity 2						
$a_0$	0.06	0.10	−0.14	0.06	0.26	13980
GDP	−0.27	0.15	−0.57	−0.27	0.03	11467
PS	0.43	0.12	0.20	0.43	0.67	11973
$\sigma_y$	0.84	0.07	0.72	0.84	0.99	13963
Conformity 3						
$a_0$	0.14	0.10	−0.05	0.14	0.33	14599
GDP	−0.20	0.14	−0.47	−0.20	0.07	12626
PS	0.43	0.12	0.20	0.43	0.65	12442
$\sigma_y$	0.87	0.07	0.75	0.86	1.00	15600
Conformity 4						
$a_0$	−0.12	0.13	−0.38	−0.12	0.14	13741
GDP	0.24	0.20	−0.16	0.23	0.63	10677
PS	−0.38	0.17	−0.71	−0.39	−0.05	10524
$\sigma_y$	0.88	0.09	0.72	0.87	1.08	13303

Supplementary Table S8: Numerical values of estimated parameters in Model 2 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $a$  represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia).  $ESS$  represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

			Quantiles			
	Mean	SD	2.5%	50%	97.5%	ESS
Individualism						
$\mu_a$	−0.14	0.23	−0.61	−0.14	0.30	7079
GDP	0.39	0.11	0.18	0.39	0.61	1914
PS	−0.28	0.13	−0.53	−0.28	−0.02	9514
$a_1$	−0.10	0.19	−0.47	−0.09	0.26	16000
$a_2$	−0.47	0.21	−0.89	−0.47	−0.03	643
$a_3$	0.26	0.17	−0.06	0.26	0.60	2562
$a_4$	−0.45	0.17	−0.78	−0.45	−0.10	512
$a_5$	−0.12	0.17	−0.45	−0.12	0.20	16000
$a_6$	0.02	0.23	−0.43	0.02	0.49	16000
$\sigma_a$	0.45	0.28	0.12	0.39	1.15	1414
$\sigma_y$	0.67	0.05	0.58	0.67	0.78	2058
Conformity 1						
$\mu_a$	0.19	0.21	−0.19	0.18	0.63	7054
GDP	−0.43	0.12	−0.66	−0.43	−0.19	11358
PS	0.43	0.13	0.17	0.44	0.68	8317
$a_1$	0.16	0.19	−0.21	0.16	0.55	11193
$a_2$	0.03	0.18	−0.34	0.04	0.37	14129
$a_3$	−0.10	0.15	−0.41	−0.10	0.19	5874
$a_4$	0.32	0.17	0.004	0.31	0.67	9951
$a_5$	0.47	0.24	0.05	0.46	0.95	5687
$a_6$	0.24	0.27	−0.27	0.22	0.82	12130
$\sigma_a$	0.37	0.25	0.09	0.32	0.99	4106
$\sigma_y$	0.63	0.05	0.54	0.63	0.75	8258

Supplementary Table S8 (cond.): Numerical values of estimated parameters in Model 2 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $a$  represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, 6: South Asia).  $ESS$  represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

			Quantiles			
	Mean	SD	2.5%	50%	97.5%	ESS
Conformity 2						
$\mu_a$	0.03	0.32	−0.62	0.04	0.66	10060
GDP	−0.24	0.15	−0.53	−0.24	0.05	13007
PS	0.60	0.17	0.28	0.59	0.92	8913
$a_1$	−0.20	0.26	−0.73	−0.19	0.31	11423
$a_2$	0.01	0.23	−0.46	0.02	0.47	16000
$a_3$	0.28	0.19	−0.08	0.28	0.65	9830
$a_4$	−0.42	0.24	−0.90	−0.42	0.05	8519
$a_5$	0.67	0.33	0.06	0.67	1.32	8361
$a_6$	−0.13	0.37	−0.89	−0.11	0.57	16000
$\sigma_a$	0.63	0.38	0.16	0.54	1.60	5735
$\sigma_y$	0.78	0.07	0.67	0.78	0.93	13591
Conformity 3						
$\mu_a$	0.23	0.45	−0.66	0.22	1.13	6600
GDP	−0.13	0.12	−0.37	−0.13	0.11	16000
PS	0.15	0.15	−0.13	0.15	0.45	9677
$a_1$	1.12	0.26	0.61	1.12	1.64	11825
$a_2$	−0.49	0.24	−0.95	−0.48	−0.03	16000
$a_3$	−0.29	0.17	−0.63	−0.29	0.04	11100
$a_4$	0.55	0.21	0.14	0.55	0.95	16000
$a_5$	0.54	0.24	0.08	0.54	1.01	16000
$a_6$	−0.05	0.38	−0.81	−0.05	0.70	16000
$\sigma_a$	0.92	0.51	0.38	0.80	2.23	6105
$\sigma_y$	0.73	0.06	0.63	0.73	0.86	16000

Supplementary Table S8 (cond.): Numerical values of estimated parameters in Model 2 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $a$  represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, 6: South Asia).  $ESS$  represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes in which 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 4						
$\mu_a$	−0.29	0.42	−1.24	−0.25	0.41	3428
GDP	0.30	0.20	−0.09	0.30	0.68	311
PS	−0.19	0.24	−0.65	−0.19	0.30	293
$a_1$	−0.28	0.32	−0.96	−0.26	0.32	5074
$a_2$	−0.61	0.35	−1.31	−0.60	0.02	797
$a_3$	0.07	0.25	−0.37	0.06	0.58	730
$a_4$	0.25	0.31	−0.30	0.24	0.86	344
$a_5$	−0.22	0.37	−1.00	−0.21	0.52	12982
$a_6$	−0.89	0.73	−2.56	−0.76	0.19	1674
$\sigma_a$	0.71	0.56	0.07	0.59	2.11	558
$\sigma_y$	0.82	0.09	0.66	0.81	1.03	2414

Supplementary Table S9: Numerical values of estimated parameters in Model 3 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $\alpha$ , GDP, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes when 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Individualism						
$\mu_a$	−0.26	0.23	−0.72	−0.26	0.22	5700
$\mu_{\text{GDP}}$	0.32	0.28	−0.26	0.32	0.86	6195
$\mu_{\text{PS}}$	−0.10	0.30	−0.64	−0.11	0.55	6511
$a_1$	−0.08	0.31	−0.61	−0.10	0.60	5328
$a_2$	−0.58	0.24	−1.06	−0.57	−0.15	5118
$a_3$	−0.01	0.20	−0.38	−0.01	0.38	4976
$a_4$	−0.49	0.19	−0.86	−0.50	−0.13	5076
$a_5$	−0.15	0.16	−0.46	−0.15	0.18	12622
$a_6$	−0.22	0.32	−0.83	−0.22	0.44	4866
GDP <sub>1</sub>	0.32	0.45	−0.61	0.32	1.26	6667
GDP <sub>2</sub>	0.32	0.21	−0.12	0.33	0.74	12025
GDP <sub>3</sub>	0.43	0.12	0.20	0.43	0.67	7035
GDP <sub>4</sub>	0.46	0.39	−0.24	0.42	1.40	6822
GDP <sub>5</sub>	0.07	0.27	−0.50	0.08	0.52	3223
GDP <sub>6</sub>	0.31	0.44	−0.61	0.32	1.21	6522
PS <sub>1</sub>	−0.37	0.25	−0.90	−0.36	0.09	6937
PS <sub>2</sub>	−0.04	0.33	−0.65	−0.06	0.63	4154
PS <sub>3</sub>	−0.50	0.17	−0.83	−0.50	−0.16	4310
PS <sub>4</sub>	0.06	0.31	−0.49	0.05	0.70	6581
PS <sub>5</sub>	0.05	0.33	−0.56	0.03	0.73	3267
PS <sub>6</sub>	0.24	0.42	−0.47	0.21	1.13	5172
$\sigma_a$	0.42	0.28	0.07	0.36	1.12	4165
$\sigma_{\text{GDP}}$	0.40	0.37	0.03	0.31	1.36	2344
$\sigma_{\text{PS}}$	0.53	0.37	0.07	0.45	1.41	2182
$\sigma_y$	0.63	0.05	0.54	0.63	0.74	6259

Supplementary Table S9 (cond.): Numerical values of estimated parameters in Model 3 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $\alpha$ , GDP, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, 6: South Asia). *ESS* represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes when 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 1						
$\mu_a$	0.28	0.22	−0.13	0.27	0.73	4268
$\mu_{\text{GDP}}$	−0.47	0.30	−1.11	−0.46	0.11	4620
$\mu_{\text{PS}}$	0.30	0.26	−0.27	0.33	0.75	4055
$a_1$	0.29	0.29	−0.24	0.27	0.91	3831
$a_2$	0.17	0.22	−0.26	0.17	0.60	4998
$a_3$	0.04	0.19	−0.34	0.04	0.40	2316
$a_4$	0.36	0.19	0.001	0.35	0.75	4749
$a_5$	0.49	0.27	0.03	0.47	1.08	2803
$a_6$	0.33	0.32	−0.26	0.31	1.03	5112
GDP <sub>1</sub>	−0.45	0.44	−1.39	−0.44	0.47	5515
GDP <sub>2</sub>	−0.58	0.24	−1.07	−0.56	−0.16	2545
GDP <sub>3</sub>	−0.36	0.13	−0.62	−0.35	−0.09	5804
GDP <sub>4</sub>	−0.42	0.39	−1.20	−0.43	0.41	3867
GDP <sub>5</sub>	−0.53	0.44	−1.51	−0.49	0.34	4718
GDP <sub>6</sub>	−0.51	0.45	−1.50	−0.47	0.37	4194
PS <sub>1</sub>	0.29	0.25	−0.23	0.30	0.73	4438
PS <sub>2</sub>	0.16	0.33	−0.58	0.20	0.69	2562
PS <sub>3</sub>	0.59	0.18	0.25	0.58	0.94	2377
PS <sub>4</sub>	0.29	0.27	−0.29	0.31	0.79	6891
PS <sub>5</sub>	0.28	0.35	−0.51	0.32	0.92	6164
PS <sub>6</sub>	0.20	0.35	−0.61	0.26	0.79	3457
$\sigma_a$	0.34	0.28	0.02	0.28	1.02	2530
$\sigma_{\text{GDP}}$	0.37	0.35	0.03	0.28	1.30	1847
$\sigma_{\text{PS}}$	0.37	0.30	0.03	0.30	1.11	1764
$\sigma_y$	0.63	0.05	0.53	0.63	0.75	11556

Supplementary Table S9 (cond.): Numerical values of estimated parameters in Model 3 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $a$ , GDP, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia).  $ESS$  represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes when 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 2						
$\mu_a$	0.04	0.35	−0.67	0.05	0.73	4806
$\mu_{\text{GDP}}$	−0.21	0.40	−0.98	−0.23	0.66	667
$\mu_{\text{PS}}$	0.60	0.27	0.08	0.58	1.14	1533
$a_1$	−0.27	0.44	−1.25	−0.27	0.52	3549
$a_2$	0.05	0.27	−0.48	0.03	0.60	2356
$a_3$	0.25	0.21	−0.15	0.24	0.69	283
$a_4$	−0.39	0.28	−0.94	−0.38	0.14	279
$a_5$	0.62	0.37	−0.06	0.59	1.39	1203
$a_6$	−0.08	0.46	−1.04	−0.06	0.86	4291
GDP <sub>1</sub>	−0.14	0.61	−1.37	−0.18	1.24	3815
GDP <sub>2</sub>	−0.13	0.28	−0.66	−0.16	0.46	2177
GDP <sub>3</sub>	−0.28	0.16	−0.59	−0.28	0.05	1805
GDP <sub>4</sub>	−0.06	0.53	−1.06	−0.12	1.27	3637
GDP <sub>5</sub>	−0.45	0.61	−1.86	−0.36	0.62	159
GDP <sub>6</sub>	−0.12	0.61	−1.31	−0.14	1.28	4112
PS <sub>1</sub>	0.71	0.31	0.20	0.66	1.41	633
PS <sub>2</sub>	0.49	0.34	−0.27	0.50	1.11	2107
PS <sub>3</sub>	0.56	0.19	0.18	0.55	0.97	1603
PS <sub>4</sub>	0.69	0.31	0.15	0.65	1.38	1173
PS <sub>5</sub>	0.58	0.38	−0.22	0.56	1.36	2393
PS <sub>6</sub>	0.58	0.35	−0.17	0.57	1.30	2213
$\sigma_a$	0.63	0.44	0.11	0.54	1.73	181
$\sigma_{\text{GDP}}$	0.58	0.55	0.07	0.40	1.86	79
$\sigma_{\text{PS}}$	0.31	0.33	0.03	0.22	1.11	275
$\sigma_y$	0.79	0.07	0.67	0.79	0.93	1738

Supplementary Table S9 (cond.): Numerical values of estimated parameters in Model 3 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $a$ , GDP, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia).  $ESS$  represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes when 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 3						
$\mu_a$	0.02	0.39	−0.73	0.01	0.79	5471
$\mu_{\text{GDP}}$	−0.56	0.55	−1.73	−0.53	0.40	6642
$\mu_{\text{PS}}$	0.20	0.26	−0.29	0.19	0.72	6302
$a_1$	0.36	0.51	−0.63	0.35	1.38	5830
$a_2$	−0.44	0.27	−0.98	−0.44	0.08	9198
$a_3$	−0.38	0.18	−0.73	−0.37	−0.04	9299
$a_4$	0.47	0.26	−0.05	0.47	0.98	7928
$a_5$	0.31	0.26	−0.20	0.30	0.82	9293
$a_6$	−0.19	0.52	−1.26	−0.17	0.82	7889
GDP <sub>1</sub>	−1.00	0.91	−3.05	−0.89	0.52	4704
GDP <sub>2</sub>	−0.69	0.29	−1.27	−0.69	−0.13	9585
GDP <sub>3</sub>	−0.04	0.13	−0.28	−0.04	0.21	11987
GDP <sub>4</sub>	−0.31	0.61	−1.53	−0.32	0.95	7879
GDP <sub>5</sub>	−1.01	0.57	−2.20	−0.98	−0.01	7017
GDP <sub>6</sub>	−0.32	0.81	−1.96	−0.33	1.40	7502
PS <sub>1</sub>	0.43	0.35	−0.12	0.38	1.22	3301
PS <sub>2</sub>	0.11	0.33	−0.62	0.12	0.76	7805
PS <sub>3</sub>	0.12	0.16	−0.20	0.12	0.43	8233
PS <sub>4</sub>	0.09	0.28	−0.54	0.10	0.62	10325
PS <sub>5</sub>	0.20	0.30	−0.39	0.19	0.84	9474
PS <sub>6</sub>	0.21	0.35	−0.49	0.19	0.95	9390
$\sigma_a$	0.71	0.44	0.21	0.61	1.80	4464
$\sigma_{\text{GDP}}$	0.88	0.66	0.16	0.72	2.54	4001
$\sigma_{\text{PS}}$	0.35	0.31	0.02	0.27	1.15	2286
$\sigma_y$	0.70	0.06	0.60	0.69	0.82	16000

Supplementary Table S9 (cond.): Numerical values of estimated parameters in Model 3 using both pathogen stress and GDP per capita as independent variables. GDP and PS represent GDP per capita and pathogen stress, respectively. Subscript numbers under  $a$ , GDP, and PS represent global regions (1: Sub-Saharan Africa, 2 East Asia and Pacific, 3: Europe and Central Asia, 4: Latin America and Caribbean, 5: Middle East and North Africa, and 6: South Asia).  $ESS$  represents effective sample sizes of MCMC simulations. Rows shaded in gray indicate the effects of slopes when 95% Bayesian credible intervals of values did not include zero.

	Mean	SD	Quantiles			ESS
			2.5%	50%	97.5%	
Conformity 4						
$\mu_a$	−0.16	0.57	−1.20	−0.11	0.67	2631
$\mu_{\text{GDP}}$	0.41	0.68	−0.86	0.37	1.89	4987
$\mu_{\text{PS}}$	−0.32	0.45	−1.32	−0.29	0.45	3228
$a_1$	−0.26	0.53	−1.38	−0.23	0.74	2023
$a_2$	−0.50	0.41	−1.33	−0.47	0.24	2997
$a_3$	0.14	0.27	−0.37	0.14	0.67	2761
$a_4$	0.23	0.37	−0.42	0.23	0.98	422
$a_5$	−0.06	0.55	−1.12	−0.07	1.06	2235
$a_6$	−0.52	0.85	−2.51	−0.36	0.79	2019
GDP <sub>1</sub>	0.55	0.93	−1.05	0.42	2.84	4174
GDP <sub>2</sub>	0.46	0.36	−0.23	0.44	1.22	4933
GDP <sub>3</sub>	0.28	0.21	−0.13	0.28	0.70	5579
GDP <sub>4</sub>	0.04	1.06	−2.69	0.21	1.82	168
GDP <sub>5</sub>	0.28	0.99	−1.95	0.32	2.21	227
GDP <sub>6</sub>	0.75	1.19	−1.03	0.52	3.81	3812
PS <sub>1</sub>	−0.05	0.43	−0.83	−0.09	0.92	1860
PS <sub>2</sub>	−0.32	0.46	−1.30	−0.31	0.58	3443
PS <sub>3</sub>	−0.13	0.27	−0.65	−0.14	0.41	3215
PS <sub>4</sub>	−0.20	0.38	−0.95	−0.21	0.55	5786
PS <sub>5</sub>	−0.60	0.80	−2.67	−0.40	0.53	3223
PS <sub>6</sub>	−0.58	0.71	−2.29	−0.42	0.52	2831
$\sigma_a$	0.72	0.69	0.05	0.59	2.21	1952
$\sigma_{\text{GDP}}$	0.83	0.90	0.04	0.55	3.15	603
$\sigma_{\text{PS}}$	0.56	0.56	0.01	0.41	2.03	857
$\sigma_y$	0.82	0.09	0.66	0.81	1.02	6635

Supplementary Tables S10: Summary of analysis results using both pathogen stress and GDP as independent variables. Parameters in parentheses indicate the parameters in Model 3. Asterisks in columns of global effects (GDP, PS,  $\mu_{GE}$ , or  $\mu_{PS}$ ) indicate that the effect was significant (i.e., the 95% Bayesian credible interval of the effect did not include zero). Numbers in columns of region-specific effects ( $GDP_j$  or  $PS_j$ ) indicate global regions in which significant effects were found (1 = Sub-Saharan Africa, 2 = East Asia and Pacific, 3 = Europe and Central Asia, 4 = Latin America and Caribbean, 5 = Middle East and North Africa, and 6 = South Asia). Rows shaded in gray indicate the best model, in which the WAIC value was smallest in three models.

Dependent variables	Model	Parameters			
		GDP	PS		
		( $\mu_{GDP}$ )	( $\mu_{PS}$ )	( $GDP_j$ )	( $PS_j$ )
<i>Individualism</i>	Model 1	*	*		
	Model 2	*	*		
	Model 3			3	3
<i>Conformity 1</i>	Model 1	*	*		
	Model 2	*	*		
	Model 3			2, 3	3
<i>Conformity 2</i>	Model 1		*		
	Model 2		*		
	Model 3		*		1, 3, 4
<i>Conformity 3</i>	Model 1		*		
	Model 2				
	Model 3			2, 5	
<i>Conformity 4</i>	Model 1		*		
	Model 2				
	Model 3				

## Supplementary Method

### Bayesian estimation of zero-order correlation coefficients

Here, we explain the method of Bayesian estimation of zero-order correlation coefficients between  $y_a$  and  $y_b$ .  $\vec{y}$  represents the two-dimensional vector of  $y$ , and  $\vec{y}$  obeys the following statistical model:

$$\begin{aligned}\vec{\mu} &= \begin{bmatrix} \mu_a \\ \mu_b \end{bmatrix}, \\ \Sigma &= \begin{pmatrix} \sigma_a^2 & \sigma_a \sigma_b \rho \\ \sigma_a \sigma_b \rho & \sigma_b^2 \end{pmatrix}, \\ \vec{y} &\sim \text{Normal}_2(\vec{\mu}, \Sigma).\end{aligned}\tag{S1}$$

$\vec{\mu}$  represents the mean vector of  $y$ .  $\mu_a$  and  $\mu_b$  represent the mean values of  $y_a$  and  $y_b$ , respectively.  $\Sigma$  represents the covariance matrix of  $y_a$  and  $y_b$ .  $\sigma_a$  and  $\sigma_b$  represent the standard deviation of  $y_a$  and  $y_b$ , respectively.  $\sigma_{ab}$  represents the covariance of  $y_a$  and  $y_b$ .  $\rho$  represents the correlation coefficients between  $y_a$  and  $y_b$ .  $\vec{y}$  obeys two variants normal distribution with two-dimensional mean vector  $\vec{\mu}$  and 2 x 2 covariance matrix  $\Sigma$ .

We estimated the parameter values of  $\vec{\mu}$ ,  $\Sigma$ , and  $\rho$  from  $y_a$  and  $y_b$  by conducting MCMC simulations. We used the uniformed distribution  $[-1, 1]$  for a prior of  $\rho$ . We calculated the 95% Bayesian credible intervals of  $\rho$ .